

SEQUENCE LISTING

<110> TANAKA, YOSHIKAZU ONO, EIICHIRO NAKAMURA, NORIKO MIZUTANI, MASAKO <120> METHOD FOR PRODUCING YELLOW FLOWER BY CONTROLLING FLAVONOID SYNTHETIC PATHWAY <130> 47237.5008/00US <140> 10/583,110 <141> 2006-06-15 <150> PCT/JP2004/019461 <151> 2004-12-17 <150> JP 2003-420046 <151> 2003-12-17 <160> 70 <170> PatentIn Ver. 3.3 <210> 1 <211> 1422 <212> DNA <213> Artificial Sequence <220> <221> CDS <222> (1)..(1371) <223> Description of Artificial Sequence: Synthetic nucleotide construct <400> 1 atg gga gaa gaa tac aag aaa aca cac aca ata gtc ttt cac act tca 48 Met Gly Glu Glu Tyr Lys Lys Thr His Thr Ile Val Phe His Thr Ser 10 gaa gaa cac ctc aac tct tca ata gcc ctt gca aag ttc ata acc aaa 96 Glu Glu His Leu Asn Ser Ser Ile Ala Leu Ala Lys Phe Ile Thr Lys 144 His His Ser Ser Ile Ser Ile Thr Ile Ile Ser Thr Ala Pro Ala Glu tot tot gaa gtg gcc aaa att att aat aat ccg tca ata act tac cgc 192 Ser Ser Glu Val Ala Lys Ile Ile Asn Asn Pro Ser Ile Thr Tyr Arg 50 ggc ctc acc gcg gta gcg ctc cct gaa aat ctc acc agt aac att aat Gly Leu Thr Ala Val Ala Leu Pro Glu Asn Leu Thr Ser Asn Ile Asn 70

65

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		_		-			_		_	-			_	atc Ile		336
														tcc Ser		384
-	_							_	_	_			_	ttt Phe		432
	_	_					_						-	cgt Arg		480
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	_	_					_							gtg Val		816
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		_	_		aaa Lys		_									1008
					ttg Leu											1056
	_			_	tcg Ser	_		_		_	_			_	_	1104
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Glu Glu His Leu Asn Ser Ser Ile Ala Leu Ala Lys Phe Ile Thr Lys 20 25 30

His His Ser Ser Ile Ser Ile Thr Ile Ile Ser Thr Ala Pro Ala Glu 35 40 45

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Ser Ser Glu Val Ala Lys Ile Ile Asn Asn Pro Ser Ile Thr Tyr Arg
50 55 60

Gly Leu Thr Ala Val Ala Leu Pro Glu Asn Leu Thr Ser Asn Ile Asn 65 70 75 80

Lys Asn Pro Val Glu Leu Phe Phe Glu Ile Pro Arg Leu Gln Asn Ala 85 90 95

Asn Leu Arg Glu Ala Leu Leu Asp Ile Ser Arg Lys Ser Asp Ile Lys
100 105 110

Ala Leu Ile Ile Asp Phe Phe Cys Asn Ala Ala Phe Glu Val Ser Thr 115 120 125

Ser Met Asn Ile Pro Thr Tyr Phe Asp Val Ser Gly Gly Ala Phe Leu 130 135 140

Leu Cys Thr Phe Leu His His Pro Thr Leu His Gln Thr Val Arg Gly 145 150 155 160

Asp Ile Ala Asp Leu Asn Asp Ser Val Glu Met Pro Gly Phe Pro Leu 165 170 175

Ile His Ser Ser Asp Leu Pro Met Ser Leu Phe Tyr Arg Lys Thr Asn 180 185 190

Val Tyr Lys His Phe Leu Asp Thr Ser Leu Asn Met Arg Lys Ser Ser 195 200 205

Gly Ile Leu Val Asn Thr Phe Val Ala Leu Glu Phe Arg Ala Lys Glu 210 215 220

Ala Leu Ser Asn Gly Leu Tyr Gly Pro Thr Pro Pro Leu Tyr Leu Leu 225 230 235 240

Ser His Thr Ile Ala Glu Pro His Asp Thr Lys Val Leu Val Asn Gln 245 250 255

His Glu Cys Leu Ser Trp Leu Asp Leu Gln Pro Ser Lys Ser Val Ile 260 265 270

Phe Leu Cys Phe Gly Arg Arg Gly Ala Phe Ser Ala Gln Gln Leu Lys 275 280. 285

Glu Ile Ala Ile Gly Leu Glu Lys Ser Gly Cys Arg Phe Leu Trp Leu 290 295 300

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aagettateg aagaactaaa getegaetgt etegttteeg acatgttett geettggaca 360
gtcgattgtg cggctaagtt cggtattccg aggttggttt tccacggaac gagcaacttt 420
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cacataacct tegteaacae egagtacate egteteegee teeteaagte etgtggeeet 180
gccgccctgg acgggctacc ggactttcgc ttcatgacta tccccgatgg cctccctttg 240
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<213> Artificial Sequence

1

<220> <223> Description of Artificial Sequence: Synthetic nucleotide construct <400> 18 cccggccgcg gccacataaa ccccatgctc aacatctgca aagccgtagc ggagaagagc 120 agccacatca acataacaat catcctaacc gaggaatggc tcggcttaat cggctcagcc 180 gacaagccgc cgaacataag ctacgccgcg ataccgaaca ttctgccgtc ggagcacgtt 240 cgcggcgagg atccacatgg tttttgggcg gctgtttggc agaagatgga ggagccggtt 300 gateggetge tggacgaget teggettaat aataacaage eggagtttgt gatageegat 360 getttettge attgggegge tgaegtggeg ggeaggagga atatteeett ggeatetgtt 420 tggccaatgt cggcgtccac gttcacggtg ctttaccact ttgaccttct cgttgaccac 480 ggacactttc cgatcgacat accagtgaat ggagatgcta ttgtggatta catcccggga 540 ctccctccag ttcgcgtcgc agattttcca aaagacataa gaaaacaaga agacgcatcc 600 ttegteetta aacteattee caacteacea aaatteatea tetteaette aatttaegae 660 ctcgaatcca agatcatcga cgctctaaag caaaaatctt ccttctcaat ctacaacatt 720 ggtcctcatg cttcctattc caaactcaaa cacatcctca actcggataa aatcacgaaa 780 cctgatcaaq ataaccccga ctacttaaaa tqqttaqatc tccaacctcc caactccqtc 840 ttgtacattt cactcggcag tttcctatcc atttccgcag cccaaatgga tgaactcgca 900 accggaatac gaaactctgg tgtccgcttt ttgtgggtgg cacgtggcga aacaaaccgg 960 ttgaaagaga tttgttgtga tcatgaaaag gggctgatca tagaatggtg cgatcaaatg 1020 caggttettt eteattette ggttggtgga ttettgtege attgtggttg gaattegaet 1080 aaagaggcgt tgatggccgg ggtgccgttt ttgactattc caattatgtt tgatcaagtg 1140 tctaacgcga aggcggtcgt ggaagattgg agggtgggt ggagggtggt gaatgagttt 1200 aatgaagaag agttggtggg aggagatgag attgcgaata ttgtgaggag gtttatggat 1260 atggaaaatg gtgagaggaa agagttgacg aaaaatgtga aagaggtgca gaagatttgt 1320 gcgagagagt tcgaagatgg agatggacag tcgtttgagt ttaatgttga aagtttggtt 1380 caattgattc tgcaattggg tccgtaa <210> 19 <211> 1428 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic nucleotide construct <400> 19 atgaacaaca caacccaaca acaaacagta gcattagcac tagcacctca ctgtttaatc 60 gtcccattcc cattccaagg ccacattaac cccttactcc aattcgccaa acgcctcata 120 actcaccaca acaaaaacct ccaaatcaca ttcgcactca ccaaattcat cctcaccaac 180 ctctcctccq qtqccqqaqa atcatccttc tctctccqqt caatctccqa cggcttcgac 240 geoggeggee gegeteagge caacteegge geogaatace tetecaaatt eegegagate 300 ggatctcaaa ccctaaccga acttatccaa gacctatccg aatcgggtcg acccgttgac 360 tgcgtggtct acgacccgtt cgtaccttgg gccttagatg ttgccaaggg taaattcgga 420 atttcaacgg cggcgttttt tacgcagtcg tgtgcggtgg ataatatata cagtcgggtt 480 tataacggcg atttggagct gccgttgccg gagaatgagg tggttagggt tccgggtttg 540 ccggagatgg agccgtttga gatgccgagc tttgtgtatt taaacgggtc gtacccgtcg 600 aqttttqaqa tqqttqtqqq tcaqtttagg aatgttgatg aggcggattg ggtttttgtc 660 aacacttttt atgagttgga gaaagaggtc attgactgga tgtcaaaatc ttggcgagtg 720 aaagcaattg gacctaccat accatcaatg ttcatggaca agagattgca agaggacaaa 780 tcatacggtc ttagcatgtt caagcataca acaaatgact gcataaattg gctcaacgga 840

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gcgatgccac agtggacgga tcagagtacg aacgctaagt ttatcgtgga tgtttggggt 1200
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tggaaggaac gggcgtgcaa tgcagttgat gaaggggga gttcagacaa aaatattgaa 1380
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<210> 20
<211> 1425
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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      nucleotide construct
<220>
<221> modified base
<222> (1359)
<223> a, t, c, g, unknown or other
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qtctcaacqc taqaqacqqc aaaqctactc gtcgatcgaa acaaacgcct caccatcaca 120
atcotcotca tgaagetgee agtegaegee aaggtagatg attectteae aaaaaatcoc 180
tectgetete aaataaettt tgtacatete eetegaateg ageacagtte catggaacca 240
ccgggaactc ccgaatcctt tgtacacagg ttcgtcgaga gccaaaaatg tctcgtaaga 300
gatgcggtgg ttaaagcaac ggagggctca aaatcaaaca ggctagccgg atttgtaatc 360
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gctttcacqt ccqqqqccqc aactctcqqq ctattqttcc atttgcagag tcttagagat 480
qaatttaatc aqqacqtqaa qgagtacgag aactcggaag ttgagatatc gatcccggct 540
tatgttaacc cgttcccttc caaatccttg ccgtctcctg tcttcaacga ggacggtgtt 600
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cccatcgggc cagtaattca cgccacggaa gataatgcaa acaaaggaaa gcaggacgaa 780
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taccggtttt tatggtcatt gagaaagccg cctcccaaag aaaaagcgga gtttccaggg 960
gagtacaaag attttaatga agttttacca gaagggttct tacaacgtac gtccgggaga 1020
ggtaaggtaa taggatgggc tccgcagatg gccgtgttgt ctcacaatgc ggtgggagga 1080
ttcgtgtcgc attgcggctg gaactcgacg ttggagagtg tttggtgcgg agtgccaatg 1140
gccgtgtggc cattggcggc cgagcaacat gcgaacgcgt tccagttggt gaaggagttg 1200
ggaattgcgg tggagattaa gatggattat aggaagaaca gtggtgtgat tgtggaggca 1260
aaaatgattg agaaaggaat cagggagttg atggacccgg aaaatgagat aaggggtaat 1320
gtgaaagtga tgaaaaagga gagtaggana gctgtcgtgg atggtgggac ttcttttgat 1380
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<210> 21
<211> 1446
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      nucleotide construct
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cacatcacat tegteaacac agagtacaac cacegeeget tgeteaagte eeteggeeee 180
gacgeteteg atggettgee ggattteega ttegeaacea teecegaegg tetteeteeg 240
totgacgogg acgtcactca ggatgttcct tototttgta tgtccaccac taacacttgc 300
ttggagccct ttaccgagtt gctgttgaaa ctcaataact ccggcccgga cgtgccaccg 360
gtgacctgca tcgtctcgga tggtgtcatg agcttcacat tgaaggcggc ggagaggttt 420
gcgctgccgg aagtgctgtt ctggacgacg agtgcgtgtg gtttcttggc gtacacgcag 480
tataagcgtc tcttggagaa aggctatgtc cctctcaaag atatgagcca gttaacaaat 540
agctatctgg aaacaaccct cgactgggtt ccaggaatga aggatatccg attaagggac 600
ttcccatcat tcatcaggac aacggatcca aaagacatca tgtacaattt cgtattacaa 660
gaaaccgacg ctgtctccag agccaaagct ctgatcatca acacctttca tacattggaa 720
cacgacgttg taaatgccct ctccaccatg tttccacgtg tttacaccat cggctctctt 780
cagetgatgt tggaccaagt teatgacaag ageettaaeg ceatcaaete caatetetgg 840
aaagaagaat cgcaatgcat cgattggctc aattcaaaag agcccgaatc cgttgtgtat 900
gtgaatttcg gtagtgtcac tgttgtgact gctcaacaac tgacggaatt tgcgtggggg 960
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teggeaatge tgeeceetga attettgaeg gacaeggaag acagaageat getaataage 1080
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ggttggaact cgacgcttga aagtattgtc agcggagtgc ctatgatatg ttggcctttc 1200
tttgctgagc aacagacaaa ttgtaggttc agttgcgtgg aatgggaaat aggaatggag 1260
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gaaaagggga agaaaatgaa gaagaaagct atggagtgga agatgaaagc agaagcagca 1380
gctgccctg ggggaccttc gtctttaaat ttggaaaaac ttattgagga ggtgcttttg 1440
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caataa
<210> 22
<211> 1308
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      nucleotide construct
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aatcacaatc gtctagtgcg tgcgagaggc cccgaatctg ttaaaggtcg cgatgatttt 180
cagttcaaaa ccatacctga tggactaccg ccttttgata aggacgcaac gcaagacata 240
cctcaactgt gtgattctct tcaaaagaat ggtcttcctc cattgttgga cctcattaaa 300
aqtattaatq attcaccqqa ctqtccaaat gttacctgta tagtgattga tttggccatg 360
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ccacttaaag atgaaagtca aataactaat ggctatcttg ataccaaact agactgggtg 540
ccagggatga agaacattag gctcagagat tttcctagtt tcatccgaac gactgatcca 600
gatgatatca tggtgaactt catgattttt aacatgaaga atgcgcctcg tgcaaaggct 660
gtggtagtca acacattcga tgaattggag aaagatgtat tggaggccct aagtaaaaaa 720
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gaggtaaaat ctataggatc aagcttgtgg aaagaagaca acacgtgcat cgcctggctc 840
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cctcaacaac tattggagtt cgcatggggc ctagccaata gcaaccatta ctttttgtgg 960
atcataaggc cagatttggt aagtggagaa tctgcgattt tatccgaaga gtactcaaag 1020
gaagttgaag ggcgggcgat gatggtgcgt tggtgctctc aagagcaagt attggcccat 1080
cetteggtag gtggattett gacacattet ggetggaact egactatega aggaatgtea 1140
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gaaggtgttc ctatgatttg ttggcctttt tttgctgacc aacagaccaa ttgtcggtat 1200
gcatgcacgg agtgggagat tggaatggag attgaaggag aggttacgag ggataaagtg 1260
gcggatttgg tgaaaatatt gatggaggag ggaaggggag agcgatga
<210> 23
<211> 1506
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     nucleotide construct
<400> 23
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atgattccca tggtagatat cgccagatta ctcgcgaagc gcggtgtcac aatcaccatt 120
ctactcacac cccacaatgc caacagggtc aaaacagtca ttgctcgtgc aatcgattca 180
ggactaaata tcaatgtcat ccacttcaaa tttccatccg ttgaggtcgg attgcccgaa 240
ggttgtgaga atttcgatat gctccctgac atcaatggcg cattgcagtt tttcaaagcc 300
actttcatgt tacaagaaca ggtcgaagag ttgcttccaa agctcgagcc tcttccgagc 360
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ccaagaattg tgtttcacgg gacaagttgc ttttctctcc tatgtatgca cgttttagga 480
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gactggacga agtttcgtga tgaggtgcga gaggctgagg taaaagcatt tggaacggtg 660
gccaatactt ttgaagattt ggaaccagag tatgtcaaag aatacagcag agttaaaggc 720
aaaa'aagtct ggtgcatagg tcctgtttca ttatgcaaca aagatggcat agacaaggcc 780
gaaaqaggta acatggcttc aatcgacgca caccattgct tgaagtggct caattcacac 840
ctgatagagc ttggattggc tttagaagca tcaaacagac cttttatttg ggtagttaga 960
gatccatcac aagaacttaa aaaatggttt ttgaatgaga aatttgagga aagggtaaag 1020
gatagaggcc ttttgatcaa cggttgggcg cctcaagtgc tcatactttc ccatccatct 1080
gttggagggt ttgtaacgca ctgcggctgg aactcgatgc ttgaaggggt tacttcaggc 1140
ttgccgatga taacgtggcc tgtatttgct gagcagtttt gtaatgaaaa gtttattgtt 1200
cacgtgatca agactgggat aagagtgggt gttgaagtgc ctatcatctt tggagatgaa 1260
gaaaaagtcg gagttttggt gaagaatgat gagataaaga tggttataga taagttgatg 1320
gatggaggag aagagggaga agagagaaga gagagagctc aaaagcttgg agaaatggca 1380
aaaaaggcaa tggaggaggg tggttcttct tatcataatt tgacatcggt catgcaagat 1440
gtcatgatgc aacaagctaa taatggagat caatatgaag atggtgttac agttataaat 1500
                                                                 1506
acatga
<210> 24
<211> 30
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      primer
<400> 24
                                                                 30
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<210> 25
<211> 36
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<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      primer
<400> 25
cccctcgagg gtacctcaca aaacattatt cacgac
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<210> 26
<211> 24
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      primer
<400> 26
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<210> 27
<211> 26
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
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<400> 27
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<210> 28
<211> 1386
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      nucleotide construct
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gaatttaacc atgaacgcct cctgagaacg agaggcccga attcccttga cgggttgcct 120
tegtttegat tegagacaat teeegaeggt etteegeeat cagaceeega tgetacacaa 180
aacgttgcat tattgtttga gtccagcaca tccaaatgct tagctccatt cagggacctt 240
cttgctaagc taaaccacac cgacgtgccg ccagttactt gcatactatc cgacttaatc 300
atgagettea etettgaage tgeteaagag eteageatee etgatgteet tttttggace 360
gctagcgctt gtggatacct cgcttatgca cactatgcca cgcttattga aaaaggattt 420
acacctttca aagatacgag ttgcttgacc aatgggtatt tggataccgt tattgatgat 480
attectagte tggaaggeat acgtetgaga gacattecaa gttttateag aacaactaat 540
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atgttgcctc cagtttacac agttggaccc ctgcatttgg ttgaaaagca tgttggtcac 720
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acaactgacc aactgattga gttttcttgg ggtcttgcta atagcaacat atccttcttg 900
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gaagccacaa aagaaagagg gtgtttagca aattggtgcc ctcaagagaa agttcttagc 1020
cacccatcca tcagaggatt cttaactcac agcggatgga attcaactct tgagagcatt 1080
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ttttgctgca caaaatgggg cataggcata gagctagaca atgatgtcaa aagggataaa 1200
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gctatggagt ggaagaagct ggccgaagag tctgcccaga gttcatcttt taagaatcta 1320
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<210> 29
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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      nucleotide construct
<400> 29
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aacgccgccg ctagttccgt cgccgacgtg gcggccatat cttatcagca actcaaaccg 180
gccactctcc cttcggatct aaccaaaaac ccaatcgagc tcttcttcga aatcccacgt 240
ctacataatc ctaacttgct cgaagcgctg gaagaactgt cactaaaatc aaaagtaagg 300
qcatttqtga tagatttctt ttgcaatccc gcatttgagg tttcgactag cttgaacata 360
cccacttact totatgtcag cagoggogg tttgggctat gcgggttctt gcattttccg 420
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ggttgcccc cggttttgtc ctcggatttt ccgaaaggta tgttctttcg caagagtaac 540
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aacgcetteg aegegatgga gtteegaget aaagaageee tegteaacaa tetgtgegta 660
cccaattcgc caactccccc agttttctta gtcggcccat tggtcggagc aagcacaact 720
acgaaaacca caaacgaaca gcacgaatgc ttgaaatggc tggacgtgca gccagacaga 780
agcgtgatct tcttatgttt cggtaggagg ggtttgttct ccgcagacca attgaaggaa 840
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ctgagccatg gcgcggttgg agggttcgtg acgcactgtg ggaggagttc gatattggaa 1080
geggtgtegt ttggggtgee gatgateggg tggeegatat aegeggagea gaggatgaat 1140
agggtgttca tggtggagga gatgaaggtg gcgttgcagt tggatgaggt ggaggaaggg 1200
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<210> 30
<211> 1362
<212> DNA
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<213> Artificial Sequence

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17 <220> <223> Description of Artificial Sequence: Synthetic nucleotide construct <400> 30 atggaggcag acaaagaaaa tctcaagatt ttaatgttcc catggttggc tcatggtcat 60 atatttccat ttcttgagct agccaaaaga atcttgaagc gaaaaaactg gcacatatac 120 ttgtgtacca cagccataaa cttcagttct atcaacaact tcattgaaaa atataagttg 180 gagaactcaa tagaagtagt agaactccat atagaaccat cccctgaact tccacctcat 240 taccacacta caaagaattt gccaacaagt ctcaattcta ccctattaaa ggccattcag 300 acgtcgaatt cgagcttctc agacatcatc agaacattga aacctgaact agtgatatat 360 gatgtgtttc aaccttgggc tgccaagatt gcttcctcac aaggtattcc tgctgtttat 420 ttttctagct ttggagggc accattatca cttatgcatc atcaccacac gtacggaaaa 480 cccgaatttc ccttccaagc aatagttgtt gaggacatcg aactggaaag tttgctctt 540 ttgtttgatt tettgtatge caacatattt gaagtggate aagattatet ttttgggaat 600 ttcaagcaat cttgtgagct tgttttgtta aagagtagta aagggattga gaggaagtac 660 atcgattatc tttcatcttt gtctcagaaa aaaatattac ctgttggacc actagtcaca 720 gttgacaata agaccaatga ggagaattcc gagatcatga attggttgag caagaaaaaa 780 caccattcaa ctgtctacat ttccttcggt agtgaatact tcctgtctaa agaagagatt 840 gaagagatag caaaagggct tgagctttgt gatgttaact ttatatggat catcagattt 900 ccagttggag tgaccgttaa cttagaagaa acactgcctc aaggtttcct tcaaagggtg 960 aacgaacggg ggatggttgt ttcaggatgg gcaccacaga gcaacatatt agcacatcca 1020 agcacaggag gctttgtgag tcactgtggg tggagttcta tcacagaaag cgtatatttt 1080 ggtgttccgg tcatagggat ggcaatgaaa cttgatcagc caataaacgc cagaatgtta 1140 tcagaggctg gtagttgtgt cgaagtcaaa agatatgaaa atgaagtgtt taggggagaa 1200 gagatagcga aggcgataaa gaaggtgatt gttgaggaca gtggagaaag gctgcggcaa 1260 agagetttag aattgagega gaagatgaaa atggaagagg aaaatgagat ggatgaagta 1320 actgagcagc tgtgggagct ttgcttgacg aaaaaacggt aa <210> 31 <211> 1437 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic nucleotide construct <400> 31 atggaacctc atatagttat attcccgttc atgtccaaag gccacacaat ccctcctc 60 cacctctccc acctcctct tagtcgcgga gtacgcgtaa cgatcttcac cactgcacaa 120 aaccaccett teategetea acatgteeca aaaacaaata atgttaccat cattgaceta 180 ccgttccctg ataacatccc tggaatttca ccaggaacgg agagcacgga caaactcccg 240 tegatgtete tettegtece gttegtgaae geegetaaat egatgeaace gttettegaa 300 gatgagettg agaaaattea tteaggggtt agttgtgtta tateggatgg ttttetteat 360 tggacgctga aatcagcatc caagttcgga attccacgac tgagtttcta cggtatgagc 420 tactatgcct tgacaatttt tcgagtcgct atctcaaaca agttaatatc attgcacgag 480 tcaccgcacg aggcattcac cttacctagt tttccttgga ttaaactcac tagagatcac 540 ttcgacaaac cacttgatca acgtgaacca aatggtccgc aatttgactt tttcatggaa 600 qcaacqacaq ctactqtqaa taqctatggt ttcttagtga atagcttcta tgagcttgaa 660 ccaactttcg cggattacta tgacaacaat tacaaaccca aggcgtggag tgtcgggcct 720 ctctgcctcg cacaaacgcc aaagaatgat aatctctcgt cgaagcctga gtggattcat 780

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<213> Homo sapiens
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<213> Artificial Sequence
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<400> 36
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acttegetag egetegagaa ategetgeat gaaaagttta taagggaega agaegagagg 180
cctaacttag catacgatca atttagcagt cagattccat tgatctctct ctctgggatc 240
gacgatgaat gtaataagag gaaagagctg tgcaagagaa tagcgcaggc atgcgaagat 300
tggggtattt ttcaagtgat cgatcatggg atcgatttga aactcgtcaa cgatatgact 360
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aatggaaggt tcaagaacgc ggatcatcga gcggtggtga attcaaacac gaatagaatg 960
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gacgacggga agcccattat agaaaagccc atcacttatg gagaaatgta caagaggaag 1080
atggctaaag acattgaact tgccaagctc aagaagctag ccaaggaaca aaagttgcaa 1140
gaagaagttg ttaataatgt tgaagatcat catcttaaca atgggaaaac taaataggag 1200
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tgtattgcct agtcggccat attatgatta aaaaaaaaa aaaaaaa
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<400> ttctct		22
<210><211><212><213>	22	
<220> <223>	Description of Artificial Sequence: Synthetic primer	
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(213)	Altificial bequence	
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010	40	
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	Artificial Sequence	
_		
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	Primer Primer	
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      primer
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<212> DNA
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					gac Asp											600
					ctt Leu											648
					gct Ala 180											696
					gga Gly											744
	-	-	_	-	gct Ala	_				_	_		_	-		792
		_		_	tac Tyr		_	_				_	_		_	840
					acc Thr											888
					gat Asp 260											936
					atg Met											984
_		_		-	aga Arg									_	_	1032
					gac Asp											1080
					att Ile											1128
				_	cat His 340		-		_				_		_	1176
					ctc Leu											1224

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gtg agg gcg aca gaa ttg gag aag cgg gtg aga gag ttg acc gag tcc Val Arg Ala Thr Glu Leu Glu Lys Arg Val Arg Glu Leu Thr Glu Ser 400 405 410	1368
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gca gag aag gcc gtg agc aag ggt gga acg tcg ctg att gca ttg gag Ala Glu Lys Ala Val Ser Lys Gly Gly Thr Ser Leu Ile Ala Leu Glu 435 440 445	1464
aaa ttc atg gac tcg att act cta taagcgtaag agttgctata aatttagcta Lys Phe Met Asp Ser Ile Thr Leu 450	1518
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tattttacaa agttacccgt attcgtttca tgtttgatac atttttcat attcgtatat	2058
gtgcccgtgt ccgtgcaata tagtaaatta gttatggtat gtgatgtttc tatgttgtaa	2118
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<210> 70

<211> 454 <212> PRT

<213> Linaria bipartita

Met Glu Asp Thr Ile Val Phe Tyr Thr Pro Ser Asp His Ser Gln Pro 5 10

Thr Ile Ala Leu Ala Lys Phe Ile Ser Lys His His Pro Ser Ile Ser 20 25 30

. **

- Met Thr Ile Ile Ser Thr Ala Ala Phe Pro Ser Ser Ala Ala Val Leu 35 40 45
- Pro Lys Thr Ile Ser Tyr His Pro Leu Pro Ala Val Pro Met Pro Pro 50 55 60
- Asn Leu Ser Ser Asn Pro Val Glu Phe Leu Phe Glu Ile Pro Arg Leu 65 70 75 80
- His Asn Thr Lys Leu Arg Glu Ala Leu Glu Arg Ile Ser Glu Thr Ser 85 90 95
- Lys Ile Lys Ala Leu Val Ile Asp Phe Phe Cys Asn Ser Ala Phe Glu 100 105 110
- Val Ser Arg Ser Leu Asn Ile Pro Thr Phe Phe Glu Ala Ser Leu Gly
 115 120 125
- Ala Ser Gly Leu Cys Glu Phe Leu Tyr His Pro Thr Phe His Lys Thr 130 135 140
- Val Pro Gly Asp Ile Ala Asp Phe Asn Asp Phe Leu Glu Ile Pro Gly 145 150 155 160
- Cys Pro Pro Leu His Ser Ala Asp Val Pro Lys Gly Leu Phe Arg Arg 165 170 175
- Lys Thr Ile Ala Tyr Lys His Phe Leu Asp Thr Ala Asn Asn Met Arg 180 185 190
- Met Ser Ser Gly Ile Leu Leu His Ala Phe Asp Ala Leu Glu Tyr Arg 195 200 205
- Ala Lys Glu Ala Leu Ser Asn Gly Leu Cys Asn Pro Asp Gly Pro Thr 210 215 220
- Pro Pro Val Tyr Phe Val Ser Pro Thr Val Ala Glu Thr Leu Ala Tyr 225 230 235 240
- Arg Glu Asn Thr Ala Ala Leu Arg His Glu Cys Leu Thr Trp Leu Asp 245 250 255
- Leu Gln Pro Asp Lys Ser Val Ile Phe Leu Cys Phe Gly Arg Arg Gly 260 265 270
- Thr Phe Ser Met Gln Gln Leu His Glu Ile Ala Val Gly Leu Glu Arg 275 280 285
- Ser Gly Arg Arg Phe Leu Trp Ala Ile Arg Ser Ser Gly Ala Gly Asn 290 295 300
- Gly Glu Pro Asp Leu Ser Val Val Leu Pro Glu Gly Phe Leu Glu Arg 305 310 315 320
- Thr Lys Asp Ile Gly Leu Val Ile Thr Trp Ala Pro Gln Lys Glu

325 330 335

Val Leu Ser His Val Ala Val Cys Gly Phe Val Thr His Cys Gly Trp 340 345 350

Asn Ser Val Leu Glu Ala Val Ser Phe Gly Val Pro Met Ile Gly Trp 355 360 365

Pro Leu Tyr Ala Glu Gln Arg Met Asn Arg Val Phe Met Val Glu Glu 370 375 380

Ile Lys Val Ala Leu Pro Leu Glu Glu Glu Ala Asp Gly Leu Val Arg 385 390 395 400

Ala Thr Glu Leu Glu Lys Arg Val Arg Glu Leu Thr Glu Ser Val Arg 405 410 415

Gly Lys Ala Val Ser Arg Arg Val Glu Glu Met Arg Leu Ser Ala Glu 420 425 430

Lys Ala Val Ser Lys Gly Gly Thr Ser Leu Ile Ala Leu Glu Lys Phe $435 \ \ 440 \ \ \ 445$

Met Asp Ser Ile Thr Leu 450